

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/602,544
Source:	17W0 -
Date Processed by STIC:	12/20/03
- 4	

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221 Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER VERSION 4.1 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- Hand Carry directly to (EFFECTIVE 12/01/03):
 U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
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Revised 10/08/03



IFWO

RAW SEQUENCE LISTING

/602 E44

DATE: 12/20/2003

Couled and Michaelle Mobides. Doed life: Combly

PATENT APPLICATION: US/10/602,544

TIME: 14:20:51

Input Set : A:\SEQLIST.txt

Output Set: N:\CRF4\12182003\J602544.raw

- 3 <110> APPLICANT: . Li-fang Liang
- 5 <120> TITLE OF INVENTION: GROWTH DIFFERENTIATION FACTOR PROMOTER AND USES
- 6 THEREFOR
- 8 <130> FILE REFERENCE: MTN-027DV1CN
- 10 <140> CURRENT APPLICATION NUMBER: 10/602,544
- 11 <141> CURRENT FILING DATE: 2003-06-23
- 13 <150> PRIOR APPLICATION NUMBER: US 09/632,879
- 14 <151> PRIOR FILING DATE: 2000-08-04
- 16 <150> PRIOR APPLICATION NUMBER: 60/092,865
- 17 <151> PRIOR FILING DATE: 1998-07-15
- 19 <150> PRIOR APPLICATION NUMBER: 60/123,270
- 20 <151> PRIOR FILING DATE: 1999-03-08
- 22 <150> PRIOR APPLICATION NUMBER: 09/354,409
- 23 <151> PRIOR FILING DATE: 1999-07-15
- 25 <160> NUMBER OF SEQ ID NOS: 9
- 27 <170> SOFTWARE: PatentIn Ver. 2.0

ERRORED SEQUENCES

- 133 <210> SEQ ID NO: 9
- 134 <211> LENGTH: 159
- 135 <212> TYPE: DNA
- 136 <213> ORGANISM: Gallus gallus
- 138 <400> SEQUENCE: 9
- 139 gaagccagga tataaatcag ttcacccttg gctgtaacca aatgctgtct agtgacttgt 60
- 140 gatcgacagg gctttaacct ctgacagcta gattcattgt tgggacaaca accaatcgtc 120
- 141 gdttttgacg acatgagcct aatcaaagtt ggagtataa 159

E--> 144 (1)

delite see p.2 for more enou

```
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<213> Gallus gallus
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      CAN BE ANY NUCLEOTIDE
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aaagctgcag tgactgtaa

VARIABLE LOCATION SUMMARY

PATENT APPLICATION: US/10/602,544

DATE: 12/20/2003 TIME: 14:20:52

Input Set : A:\SEQLIST.txt

Output Set: N:\CRF4\12182003\J602544.raw

Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing. Use of <220> to <223> is MANDATORY if n's or Xaa's are present. in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:5; N Pos. 9,30,32,50,55,92,114,146,149,151,154,158,161

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/602,544

DATE: 12/20/2003 TIME: 14:20:52

Input Set : A:\SEQLIST.txt

Output Set: N:\CRF4\12182003\J602544.raw

L:88 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:5 L:88 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:5 L:88 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0

M:341 Repeated in SeqNo=5

L:144 M:254 E: No. of Bases conflict, this line has no nucleotides.